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## CDS: SPBC16G5.09



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Comparison of the Comparison	Term
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	soelectric point pH 4.5 Charge -24.5
socieeti le politi pri 115 Charge 2115	sociecti ic point pri 4.5 Chai ge -24.5

Transmembrane

21 and 22

1 probable transmembrane helix predicted for SPBC16G5.09 by TMHMM2.0 at aa 469-491

Other genes

annotated to this

228 others

Gene Ontology Annotation

IEA (GOA:interprolGO REF:0000002) with

integral to	membrane	IEA (GOA:spkwlGO_REF:0000004) with SP_KW:KW-0812	793 others
trans-Golgi network		ISS (PMID:17072883) with SGD:S000003171	2 others
vacuole		IDA (PMID:16823372)	145 others
Molecular carboxyper activity		ISS (PMID:17072883) with SGD:S000003171	none
		Catalytic Activity	
EC 3.4.16.0	6: IUBMB		
		Published Expression Profiles	
Gene Expre	ession Viewer	r Cell Cycle Meiosis Environmental Stress Pheromone Response/Mating	
Transcripto	meViewer	SPBC16G5.09 High-resolution view of transcripts in neighbourhood	
		Literature	
Search for i	in PubMed		
	•••••	Domain Information	
View Pfam	domain struc	cture for this gene product	
View SCOI	P superfamily	!	
DB	Accs	Description	
Pfam	PF00450	Serine carboxypeptidase	
MEROPS	S10.007	MEROPS	
InterPro	IPR001563	Peptidase S10, serine carboxypeptidase	
PRINTS	PR00724	Carboxypeptidase C serine protease (S10) family signature	
ProDom	PD001189	ProDom	

Database Cross-References

Schizosaccharomyces pombe chromosome II

SPBC16G5.09 Fission yeast strain database, National BioResource Project (Japan)

http://www.genedb.org/genedb/Dispatcher?formType=navBar&organism=pombe&desc=yes&wildcard=ye...

Carboxypeptidase kex1 precursor (EC 3.4.16.6) (Carboxypeptidase D).

Domains

GPI Anchor

proteolysis

DB

**EMBL** 

NCBI

NBRP

YOGY

PIR

Biotwiki

Entrez Gene FYSSION

Accs

CU329671

T39601

SPBC16G5 09 Biotwiki

FYSSION SPBC16G5.09 FYSSION GermOnline SPBC16G5.09 GermOnline

SPBC16G5.09 NCBI Entrez Gene

PIR

SPBC16G5.09 Retrieval of eukaryotic orthologs

UniProtKB O60123

Description

Term (browse Amigo)

Biological Process

Cellular Component

Not found

Qualifier Evidence

InterPro:IPR001563

UniProtKB Annotation For This Protein

Catalytic Activity Preferential release of a C-terminal arginine or lysine residue.

Function Protease with a carboxypeptidase B-like function involved in killer toxin and alpha-factor precursor processing (By similarity).

Similarity Belongs to the peptidase S10 family.

Subcellular Location Vacuole membrane; Single-pass type I membrane protein.

Carboxypeptidase (5 others), Complete proteome (4982 others), Glycoprotein

Keywords (169 others), Hydrolase (451 others), Membrane (801 others), Protease (90 others),

Signal (136 others), Transmembrane (651 others), Vacuole (42 others)

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